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Radiocarbon Content of CO₂ Respired from High Arctic Tundra in Northwest Greenland

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Abstract

Tundra soils contain large amounts of organic carbon (C) that might become available to microbial decomposition as soils warm. To elucidate the C sources currently sustaining CO₂ emissions from striped tundra soils (soil respiration) in Northwest Greenland, we studied the seasonal pattern and radiocarbon (¹⁴C) signature of soil respiration and of CO₂ within the pore space, respired from roots and non-root–associated microbes, and of bulk soil organic matter.

Old C pools are present in the topsoil of both barren ridges (1000–5000 yrs) and vegetated troughs (modern to 600 yrs). Before leaf-out, soil respiration was depleted in ¹⁴C relative to atmospheric CO₂, root and microbial respiration within the topsoil, demonstrating a substantial contribution of C fixed before 1950. As the growing season progressed, the contribution of older C pools decreased, but remained apparent in the soil respiration from ridges and in pore space CO₂. Soil respiration from troughs became dominated by recently fixed C.

As the active layer deepens with permafrost thaw, buried C may become an increasingly larger component of soil respiration. Detecting microbial decomposition of older C pools requires continuous monitoring of soil and microbial respiration and better constraints on soil C pools.

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Introduction

Soils of the arctic permafrost region contain an estimated 496 Pg organic carbon (C) within the top meter and 1024 Pg C within the top three meters (Tarnocai et al., 2009). Recent climate change (Box, 2002; Hinzman et al., 2005; Lawrence et al., 2008b) may intensify microbial activity (Schimel et al., 2006) and amplify losses of C to the atmosphere as carbon dioxide (CO_2) (Jones et al., 2000; Nordstroem et al., 2001; Illeris et al., 2003; Welker et al., 2004) or methane (Mastepanov et al., 2008). Increased C losses could turn the region from a C sink to a source providing a positive climate feedback until increases in plant productivity and vegetative cover offset respiratory losses (Oechel et al., 1995; Sullivan et al., 2008a; Schuur et al., 2009).

The High Arctic's (N of $\sim 70^{\circ}$ latitude) C biogeochemistry is poorly constrained by measurements, especially for prostrateshrub tundras (539 $\times 10^{6}$ km²; Walker et al., 2005) where patterned ground (Kessler and Werner, 2003) complicates spatially accurate estimates of soil C pools and fluxes (Michaelson et al., 1996). A recent study (Horwath, 2007; Horwath et al., 2008) concluded that high arctic soils (1.984 $\times 10^{6}$ km²) contain 12.06 Pg C—an order of magnitude more C than previously estimated (Bliss and Matveyeva, 1992). Given that permafrost thawing will expose deeper-lying pools of soil C, the lack of data on C depth distributions and its exchange with the modern atmosphere constrains our ability to model accurately the panarctic C cycle and its climate feedback potential (Sitch et al., 2007).

The main pathway returning C from well-drained arctic soils to the atmosphere is decomposition of organic matter by microorganisms to CO_2 (soil respiration). There are multiple sources of soil C, including above- and belowground plant litter,

root exudates, and microbial byproducts that may vary in age from days to millennia (Horwath et al., 2008). Quantifying the relative contribution of these multiple sources to the total respiration flux is especially important today as the deepening of the active layer with climate warming can result in older, previously inactive soil C becoming a larger component of the modern ecosystem respiration flux (Schuur et al., 2009).

One technique to assess contributions of different sources of C to the overall CO₂ fluxes from ecosystems involves radiocarbon (¹⁴C) measurements of soil respiration and its main sources (Trumbore, 2006). This approach quantifies the ¹⁴C content of CO2 respired from plant roots and microorganisms whose C sources are plant exudates or short-lived roots (root and rhizosphere respiration) and from microorganisms decomposing other organic matter (microbial respiration). The ¹⁴C content of respired CO₂ is a measure of the time interval since this C was fixed by plants from the atmosphere. The technique can be used to estimate the mean age of C pools in soils on several time scales. Radiocarbon dating measures the mean age of C in pools that are hundreds up to about 50,000 yrs old-based on radioactive decay after an organism dies. In addition, we can use 'bomb'-C, produced by aboveground nuclear weapons testing in the 1950s and 1960s as a tracer to estimate the mean age of soil C pools that cycle on decadal to centennial time scales. Here, the ¹⁴C content of a C pool or flux is compared to the ¹⁴C content of atmospheric CO₂ between 1950 and the year of sampling. The ¹⁴C content of CO₂, and also that of recent photosynthetic assimilates, peaked in the northern hemisphere in 1963-1964 and is declining as this 'bomb' ¹⁴CO₂ exchanges with C reservoirs in the oceans and biosphere, and becomes diluted by fossil-fuel derived (>50,000 yrs) CO₂ (Levin et al., 2008).

 TABLE 1

 Overview of sampling sites (from Horwath, 2007).

		South Mountain (Akînarssuaq) #79-2004	Polar Desert #34-2005	
Latitude/Longitude	N/W	76°30′37″/68°40′14″	76°25′38″/68°59′01″	
Elevation	m	213	338	
Slope	%	3	9 to 10	
Aspect	o	180	180	
Ground pattern	_	Non-sorted stripes	Non-sorted stripes	
Soil type	_	Cryosol	Cryosol	
Active layer depth	m	0.61 to >1.2	0.97	
Total organic carbon	$kg m^{-2}$	9.44	7.16	
Organic carbon	%-dw	0.14 to 6.25	0.12 to 5.73	
Total nitrogen	%-dw	0.01 to 0.26	0.00 to 0.39	
Soil pH	_	7.37 to 9.25	7.55 to 8.42	
Bulk density	g cm ⁻³	1.07 to 1.98	0.99 to 1.82	
Vegetation ¹	_	(high arctic) prostrate dwarf shrub tundra (polar semi-desert)		
Cover	% area	37 (30 vascular)	50 (45 vascular)	
Dominant species	_	Dryas integrifolia VAHL		

¹ Vegetation is only present in troughs; ridges are mostly barren.

Our study focused on two questions: (1) How much do root and microbial respiration contribute to the overall soil respiration flux in high arctic polar semi-deserts? (2) Are soils currently losing old C that had been unavailable to microorganisms under the climatic conditions of the last few centuries?

Material and Methods

SITE DESCRIPTION

We studied two high arctic prostrate dwarf shrub tundra (or "polar semi-desert") systems ("South Mountain" and "Polar Desert") in Northwest Greenland near Thule Air Base (76°32'N, 68°50'W; 200-350 m a.s.l.) (Table 1) as part of the U.S. National Science Foundation Biocomplexity Program (Walker et al., 2008). Mean annual air temperature averages -11.6 °C, with a mean annual precipitation of 112 mm (1971-2004) (Thule Air Base unpublished; from Sullivan et al., 2008b). During the growing season (June, July, August) air temperature averages 3.5 °C. Annual air temperature has increased by 0.5 °C per decade from 1971 to 2000 (Sullivan et al., 2008b). The soils are Cryosols with an active layer depth of about 0.6 to 1.2 m (Table 1). The underlying geology is composed of Proterozoic dolostone, siltstone, and sandstone covered by glacial drift of mixed lithology (in Horwath et al., 2008). Due to a gentle slope (<10%), the soil surface is dominated by nonsorted stripes (Fig. 1): alternating 0.5-m-wide, vegetated troughs and 1- to 3-m-wide, barren ridges. Polar stripes form from frost cracking, eolian in-filling, colonization by higher plants, and freeze/ thaw dynamics. This landscape type represents $\sim 1/3$ of the entire high arctic terrestrial land cover (Bliss and Matveyeva, 1992).

SOIL RESPIRATION RATES AND PORE SPACE CO₂ CONCENTRATIONS

We measured soil respiration and the CO₂ concentrations in the soil pore space as well as in ambient air approximately weekly throughout the growing season of 2007. Soil respiration was measured using paired, dynamic chambers (30 cm i.d., \sim 8 L air V) in troughs (n = 3) and ridges (n = 3) at both study sites. To restrict diffusion of ambient air into the chambers, chamber bases were inserted about 5 cm into the mineral soil in late May 2007, sealed with soil material on the outside, and left in place until the end of the growing season. Vegetation was not clipped. Respiration rates were measured by closing a chamber with an opaque lid and circulating the air in the chamber's headspace between the chamber and an infrared gas analyzer (LI-6200, LI-COR Biosciences, Lincoln, Nebraska, U.S.A.). The respiration rate was calculated from the slope of time vs. CO_2 concentration curves (Gaudinski et al., 2000). With each chamber measurement we recorded air and soil temperature at 5 cm depth.

At South Mountain only, we measured the concentration of CO_2 in the soil pore space. Stainless steel probes (0.35 cm i.d., 0.6 cm o.d.) were inserted vertically into the mineral soil to 20, 30, or 60 cm depths in ridges and to 20 cm in troughs, and all were closed off with a rubber septum (Blue septa, Grace, Deerfield, Illinois, U.S.A.). Probes were inserted as soon as the thaw of the active layer allowed and remained in the same location until the end of the growing season. Soil gas was obtained with 60 mL syringes (BD, Franklin Lakes, New Jersey, U.S.A.). After discarding the air present within the probe, 60 mL of soil air was injected into an infrared gas analyzer (LI-800, LI-COR) and the concentration monitored with a data logger (LI-1400, LI-COR).



FIGURE 1. Picture of non-sorted, polar stripes in NW Greenland. Stripes consist of alternating 0.5-m-wide, vegetated (mountainavens [*Dryas integrifolia* VAHL] and arctic willow [*Salix arctica* PALL.]) troughs and 1- to 3-m-wide, barren ridges.

CO2 SAMPLING FOR ISOTOPE ANALYSIS

Isotope samples were taken once a month. To sample CO₂ from soil respiration, CO₂ from ambient air was removed from the flux chambers by circulating the air in the chambers' headspace through a CO₂ trap (soda lime, Fisher Scientific, Pittsburgh, Pennsylvania, U.S.A.) for 25 min at a flow rate of ~1 L min⁻¹ (>3 × chamber V). Afterwards the chamber was left closed for approximately 24 hrs to accumulate sufficient CO₂ for ¹⁴C analysis (~0.5–1 mg C). The CO₂ was sampled by circulating the head space air through drierite (W.A. Hammond Drierite Co. Ltd., Xenia, Ohio, U.S.A.) followed by an activated molecular sieve trap for 15 min (Gaudinski et al., 2000). The molecular sieve (powder-free 13× 8/12 beads, Grace) was pre-conditioned prior to the construction of a trap by baking at ≤750 °C under vacuum for ≥12 hrs. Traps were activated by baking at 650 °C under vacuum for ≥45 min (see below).

We obtained air from within the soil pore space by connecting 2 L evacuated stainless steel canisters to the soil probes. To minimize the disturbance of the soil CO₂ concentration gradient and the risk for sampling soil air from other than the sampling depth, the cans were filled via stainless steel capillaries (0.010 \times 0.063 \times 30 cm, Fisher Scientific), which restrict the flow rate, over an approximately 4 hour period (Gaudinski et al., 2000). CO₂ from ambient air was sampled by flushing air through drierite followed by a molecular sieve trap for 15 min.

To sample CO₂ respired from roots we manually extracted all roots from a block of 0–5 cm mineral soil the same day the soil was sampled. Roots were rinsed with water and placed into a 1 L mason jar. The air in the jar was circulated through soda lime for 10 min at a flow rate of ~1 L min⁻¹ to remove CO₂ from ambient air. After 12 hrs, the evolved CO₂ was collected on a molecular sieve trap for 10 min (~1 L min⁻¹).

To sample CO2 respired by microorganisms we collected soil cores (0–10 cm mineral soil) from troughs (n = 6) and ridges (n =6) in early June 2007 using a steel tube (approximately 5 cm i.d.). Sampling the surface soil does not capture the respiration of all microorganisms in the active layer, but it should provide an estimate of the dominant fraction of microbial respiration, since microbial activity in soils declines with depth, often in parallel with the distribution of roots. To ensure structural integrity, samples were wrapped in aluminum foil and packed into plastic bags in the field. Samples were frozen and transported to the University of California (UC) Irvine on ice. The frozen cores were placed into 1 L mason jars. To conserve their structural integrity, the cores remained in the aluminum foil, but the foil was removed from the top and pierced along the core to facilitate gas and water flow. Soils were thawed at 6.4 °C for four days. Before and after thawing, jars were flushed with CO2-free air (ambient air flushed through soda lime). Two out of the six cores from each site and landscape position (ridge or trough) remained at 6.4 °C, two were moved to about -4.7 °C, and two to 15.0 °C. After two weeks, the headspace air was transferred to evacuated 0.5 L canisters. Throughout the incubation period CO₂ concentrations were monitored every 1-5 days (data not shown); CO₂ concentrations in the jars were kept $\leq 2\%$).

CARBON ISOTOPE ANALYSES OF CO2

Carbon dioxide was released from molecular sieve traps by baking at 650 °C for \geq 45 min or extracted from canisters using a vacuum line, purified cryogenically and reduced to graphite using Zn reduction (Xu et al., 2007). A split of the CO₂ was analyzed for its stable C isotope ratio (DeltaPlus, GasBench II, Thermo Scientific, Waltham, Massachusetts, U.S.A.). The ¹⁴C content of the graphite was measured with accelerator mass spectrometry (NEC 0.5MV 1.5SDH-2 AMS) at the KCCAMS laboratory of UC Irvine (Southon and Santos, 2007). Radiocarbon data are reported relative to NIST OX-I (SRM 4990a) and OX-II (SRM 4990c) following Stuiver and Polach (1977).

Atmospheric CO₂ was removed from the chambers prior to sampling (see above), however, leakage of atmospheric CO₂ into the chambers during the accumulation period cannot be excluded in these rocky soils. Leakage can affect the ¹⁴C content of the sampled CO₂, in particular when respiration fluxes are low. We calculated the amount of ambient CO₂ in each sample (f_{air}) based on the δ^{13} C ratios of CO₂ in the sample ($\delta^{13}C_{obs.}$), in ambient air ($\delta^{13}C_{air}$), and respired from either roots or microbes ($\delta^{13}C_{resp.}$) (Fig. 4c):

$$f_{air} = \frac{\left(\delta^{13} C_{obs.}\right) - \left(\delta^{13} C_{resp.}\right)}{\left(\delta^{13} C_{air}\right) - \left(\delta^{13} C_{resp.}\right)} \tag{1}$$

The calculation of f_{air} is sensitive to the choice of $\delta^{13}C_{resp.}$. We estimated f_{air} for each soil respiration sample using the δ^{13} C of root respiration (-28% in June, -26% in July and August). In addition, we also calculated f_{air} using the minimum and maximum of observed δ^{13} C of microbial respiration (-19 or -29%).

Using these three estimates for f_{air} , we then calculated three air-corrected ¹⁴C signatures ($\Delta^{14}C_{cor.}$) from each observed soil respiration ¹⁴C signature ($\Delta^{14}C_{obs.}$):

$$\Delta^{14}C_{cor.} = \frac{\left(\Delta^{14}C_{obs.} - \left(f_{air} \times \Delta^{14}C_{air}\right)\right)}{\left(1 - f_{air}\right)} \tag{2}$$

We report air-corrected Δ^{14} C signatures of soil respiration as average \pm SD (n = 3). The measurement uncertainty for Δ^{14} C was <2‰.

SOIL ORGANIC MATTER AND CARBONATE ANALYSIS

After incubation, soils were dried at 60 °C and sieved to <2 mm. A subsample was ground and carbonates were removed by shaking overnight with 0.5 M HCl. Rinsed, freeze-dried samples were analyzed for C content and stable isotopic composition in duplicate (Carlo Erba, Milano, Italy; DeltaPlus, Thermo Scientific). For each site and landscape position, the samples with the highest and lowest C content were analyzed for their ¹⁴C content. Samples were combusted to CO₂ in precombusted, evacuated quartz tubes (6 mm i.d.) with CuO and Ag for 2 hrs at 900 °C, and the CO₂ was processed as described above.

For each site, we analyzed other subsamples (ridges, n = 2; troughs, n = 1) for their concentration and isotopic composition of inorganic C (carbonates). CO₂ was released from carbonates by acidifying soil samples (0.1 g trough, 1 g ridge soils) with 0.8 mL phosphoric acid in pre-evacuated glass tubes and purified on a vacuum line and graphitized as described above.

Results and Discussion

SOIL CARBON POOLS

The concentration of organic C in the mineral soil (0–10 cm) was significantly higher in troughs than in ridges, averaging 2.8 ± 0.3 vs. $0.8 \pm 0.2\%$ C (mean \pm SE, n = 12; *t*-test p < 0.01) (Table 2). Concentrations in troughs or ridges were similar at both

TABLE 2

Concentration and isotopic properties of organic matter and carbonates in the top 10 cm of mineral soil (<2 mm). Organic matter data are averages (SD), carbonate data are point samples.

	Organic carbon				Carbonates			
	Concentration (%-dw)	δ ¹³ C (‰)	Δ^{14} C (‰)	Conv. ¹⁴ C age (yrs)	Concentration (%-dw)	δ ¹³ C (‰)	Δ ¹⁴ C (‰) Conv	v. ¹⁴ C age (yrs)
South Mountain								
Ridges	1.12 (0.89)	-25.1 (0.2)	-332.0	3185	≤0.01	-	-	-
Troughs	3.18 (1.17)	-26.2 (0.6)	-24.1	140	< 0.01	-	-	-
			-81.9	630				
Polar Desert								
Ridges	0.53 (0.51)	-25.0 (1.8)	-147.9	1230	0.09	2.1	-890	17,710
			-280.2	2585				
			-469.7	5040				
Troughs	2.43 (0.88)	-26.7 (0.5)	21.5	Modern	0.16	-	-901	18,490
			9.8	Modern				
			-52.1	375				
n	6	6	2 to 3	1 to 3	1 to 2	1	1	1

sites and in the same range as previously reported (Horwath, 2007; Horwath et al., 2008).

As expected, the δ^{13} C signature of the soil organic C indicated that C₃ plants were the dominant source (-25.7 ± 0.2‰; Table 2). The ¹⁴C signature of organic C in troughs was younger than in ridges (Table 2). Radiocarbon ages in troughs or ridges were similar at both sites, although spatial variability was very high. The mean conventional ¹⁴C ages of C varied from 630 yrs to modern in troughs, and from 5040 to 1230 yrs in ridges.

The temporal and spatial dynamics of cryoturbation in striped soils are poorly understood (Walker et al., 2008). Nevertheless, the distribution of organic C and its ¹⁴C ages within the surface mineral soil of these landscapes seems to reflect the current distribution of vegetative cover. The surface soil of vegetated troughs receives annual inputs of plant leaf and root litter. In the barren ridges, organic matter is either a remnant of past vegetative cover and/or has and is being transported from troughs via cryoturbation and as dissolved organic matter. The first detailed mapping of age vs. depth profiles by Horwath et al. (2008) showed that at South Mountain troughs are associated with more stable, sand-rich A horizon wedges and ridges with more mobile, silty loam and carbonate-cemented C horizons. In troughs, ¹⁴C ages of bulk soil organic matter increased with depth (0-60 cm) from 65 to 2695 yrs, while ridges contained pockets of buried organic matter between 50 and 70 cm depth with ¹⁴C ages of about 30,000 yrs.

The carbonate content of the soils was low (<0.2%-mass) and not all samples yielded enough CO₂ for isotope analysis (Table 2). Carbonates had ¹⁴C ages older than 17,000 yrs and a δ^{13} C ratio of +2%. Thus, carbonate Δ^{14} C signatures were significantly depleted and carbonate δ^{13} C ratios were significantly enriched relative to the isotopic signatures of soil organic C.

SOIL PORE SPACE CO₂

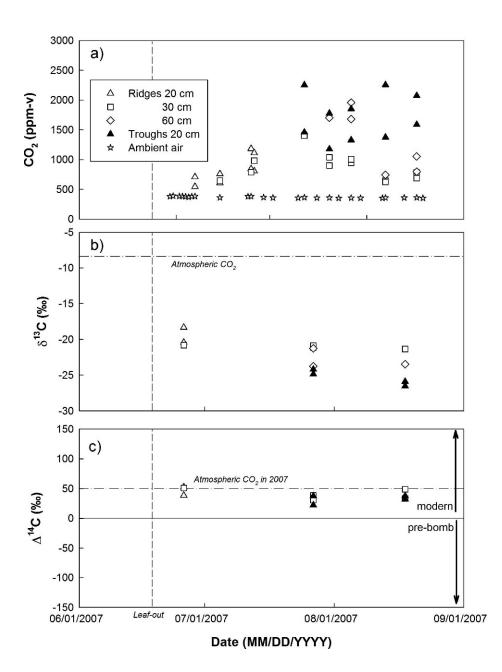
Concentrations of CO₂ in the soil pore space (measured only at South Mountain) were consistently higher in troughs than ridges for a given soil depth, averaging 1713 \pm 76 vs. 752 \pm 76 ppm CO₂ at 20 cm depth (mean \pm SE, n = 5-6, *t*-test p < 0.01) (Fig. 2a). The higher CO₂ concentrations in troughs are a result of higher root abundance and the higher microbial activity sustained by root exudates (Illeris et al., 2003) along with the overall larger amounts of organic matter. In ridges, pore space CO₂ concentrations increased with depth (892 \pm 89 ppm at 30 cm [n = 8], 1298 \pm 271 ppm at 60 cm [n = 4]), and exhibited a seasonal pattern with a maximum in mid-summer (1404 ppm at 30 and 1819 ppm at 60 cm), when soil temperature peaked (Fig. 2a).

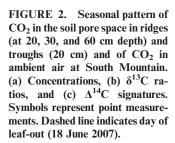
The ¹⁴C signature of pore space CO₂ throughout the growing season was modern (1950 to 2007, $\Delta^{14}C \ge 0\%$), but similar or depleted relative to the ¹⁴C signature of CO₂ in ambient air (Fig. 2c). This suggests that the respired CO₂ was a mixture of C fixed during the bomb-period (1950 to 2007, $\Delta^{14}C \ge 0\%$) and older C (pre-1950, Δ^{14} C < 0‰). The ¹⁴C signature of pore space CO2 reached a minimum in mid-summer when CO2 concentrations and soil temperatures peaked. This decrease in ¹⁴C signature was not related to changes in the proportion of leakage of ambient CO_2 into the wells: The $\delta^{13}C$ signature of pore space CO_2 at a given depth remained constant over time (Fig. 2b). Instead, the decrease implies that the contribution of older C was largest during mid-summer. Measurements of CO₂ in the soil pore space provide evidence that older C pools are being decomposed, and that decomposition rates increase as a function of soil temperature. It is unlikely that carbonates were the source of the observed ¹⁴C-depleted soil respiration flux, because carbonate concentrations were low, and a substantial contribution of carbonatederived CO₂ is not apparent in the δ^{13} C ratios of pore space CO₂ (Fig. 2b) or soil respiration (Fig. 4c).

SOIL RESPIRATION

At both sites, soil respiration fluxes were higher from troughs than ridges (Figs. 3a, 3b). In troughs, soil respiration showed a strong seasonal pattern, with a maximum in mid-summer when air and soil temperatures peaked (Figs. 3c, 3d), especially at Polar Desert (Fig. 3b). On the ridges, respiration fluxes were relatively constant over time (Figs. 3a, 3b). Soil respiration was positively correlated with soil temperature (Figs. 3e, 3f), except for ridges at South Mountain. The slightly lower respiration flux from the troughs of South Mountain compared to Polar Desert is likely related to lower plant abundance (Table 1).

Before leaf-out (18 June 2007, day 168), soil respiration rates in troughs and ridges rapidly increased in response to thawing events (= snow melt) and decreased during subsequent colder periods (Figs. 3a–3d). The most likely explanation for these repeated large, but short soil respiration events is that warming and the associated increase in soil moisture triggered the growth of microbes, while subsequent freezing and decline in soil moisture sharply reduced their activity. Microbial biomass turnover associated with freeze-thaw cycles may account for a significant





proportion of annual soil respiration (Schimel and Clein, 1996; Wallenstein et al., 2007). In contrast, it is unlikely that we observed the release of stored CO₂ that had accumulated in soils during the winter. The snow pack at our site is thin (less than 25 cm; M. Rogers, personal communication, 2008) and biological activity is restricted by very cold soil temperatures (<-10 °C) and therefore low soil moisture—limiting both the production of CO₂ and the formation of diffusive barriers such as ice lenses.

Soil respiration in troughs had modern ¹⁴C signatures (Fig. 4a). Before leaf-out, soil respiration tended to be depleted in ¹⁴C relative to the ¹⁴C signatures of ambient CO₂ as well as root and microbial respiration at 0–10 cm depth. As for soil pore pace CO₂, these signatures between 0‰ and 50‰ indicate that the respired CO₂ was a mixture of modern (1950 to 2007; $\Delta^{14}C \ge 0\%$) and older C (pre-1950; $\Delta^{14}C < 0\%$). After leaf-out, ¹⁴C signatures of soil respiration were similar to those of CO₂ in ambient air, root, and microbial respiration.

In ridges, the ¹⁴C signatures of soil respiration showed a similar seasonal pattern as in troughs, with more depleted signatures at the beginning of the growing season and more

enriched signatures later in the season (Fig. 4b). In comparison 14 C signatures of soil respiration in ridges were more variable than in troughs. Also, ridge soil respiration originated from older C sources than in troughs—indicated both by depleted 14 C signatures (pre-bomb C or mixture of bomb and pre-bomb C) and enriched 14 C signatures (bomb C) relative to that of ambient CO₂—and losses of older C could be detected throughout the entire growing season.

HOW MUCH DO ROOT AND MICROBIAL RESPIRATION CONTRIBUTE TO SOIL RESPIRATION?

Root-respired CO₂ had a ¹⁴C signature of 58.8 \pm 4.3‰ (Mean \pm SD, n = 6) and there was no difference between the two sites (*t*-test, p = 0.90) (Fig. 4a). The ¹⁴C signature of root respiration was thus enriched in ¹⁴C relative to CO₂ in ambient air (50 \pm 1‰; n = 7) by 4–14‰ (Fig. 4a). Assuming that the ¹⁴C signature of CO₂ in ambient air has been recently declining at a rate of 5.5‰ per year (Levin et al., 2008), roots were respiring C that was fixed

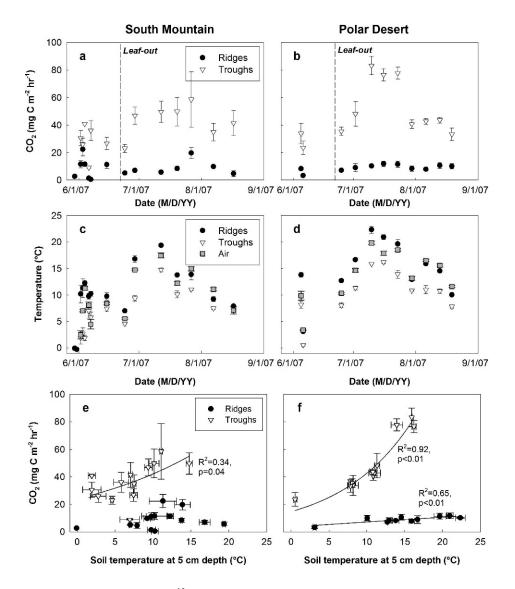


FIGURE 3. Seasonal pattern of soil respiration (panels a and b) from the surface of troughs and ridges and of air and soil temperature (panels c and d) at South Mountain and Polar Desert in 2007. Dashed line indicates day of leaf-out. Panels e and f show the correlation of soil respiration to soil temperature at 5 cm depth.

on average 1–3 years ago. The δ^{13} C ratio of root respiration was typical for C₃ vegetation (-27‰) (Fig. 4c).

The ¹⁴C signature of microbial respiration from a given soil sample was similar during thawing and subsequent incubation at a range of temperatures (data not shown). Therefore, data was analyzed after averaging ¹⁴C signatures obtained during and after thawing. This insensitivity of ¹⁴C signatures to temperature during short-term incubations confirms earlier findings (Czimczik and Trumbore, 2007).

The ¹⁴C signatures of microbial respiration exhibited large spatial variability (Fig. 4). In troughs, microbial respiration had a ¹⁴C signature of 74 \pm 13‰ (Mean \pm SD, n = 12), with no difference between sites (*t*-test, p = 0.08) (Fig. 4a). Microbial respiration from troughs was thus enriched in ¹⁴C relative to root respiration and to CO₂ in ambient air, suggesting that this C was on average older than the C respired by roots, but fixed after 1950 ($\Delta^{14}C \ge 0\% =$ modern).

In ridges, the mean ¹⁴C signature of microbial respiration was $34 \pm 9\%$ at South Mountain and $54 \pm 7\%$ at Polar Desert (Mean \pm SD, n = 4 [SMnt.] or 6 [PD]) (Fig. 4b). This variability between locations probably reflects the observed high spatial variability of soil organic C pools (Horwath et al., 2008) rather than a difference in C turnover. Microbial respiration from ridges at South Mountain was depleted in ¹⁴C relative to ambient CO₂, but still modern—indicating that microorganisms were decomposing a

mixture of recently fixed and older (pre-bomb) C. At Polar Desert, ¹⁴C signatures slightly enriched relative to ambient CO₂ suggest that microorganisms were decomposing a mixture of modern C sources. The δ^{13} C ratio of microbial respiration ranged from –19 to –29‰ (Fig. 4c).

A clear-cut partitioning of the net soil respiration flux into its sources (root and microbial respiration at 0–10 cm depth) was not feasible in troughs or ridges, because the ¹⁴C signature of soil respiration was either lower (or higher) than the ¹⁴C signature of both root and microbial respiration or there was no clear difference between the ¹⁴C signatures of the two sources (Figs. 4a, 4b). Clearly, future soil respiration partitioning approaches need to consider organic C pools within the entire active layer as potential sources for microbial respiration, and additional measurements are needed to better constrain the spatial variability of organic C sources.

ARE SOILS CURRENTLY LOSING OLD C?

Pore space ${}^{14}\text{CO}_2$ measurements (made after leaf-out) indicate that decomposition of older C intensified with increasing soil temperature and a deepening active layer and peaked in midsummer. In the soil respiration flux, however, losses of older C were most pronounced early in the season (before leaf-out) and became obscured by larger fluxes from plant roots and/or

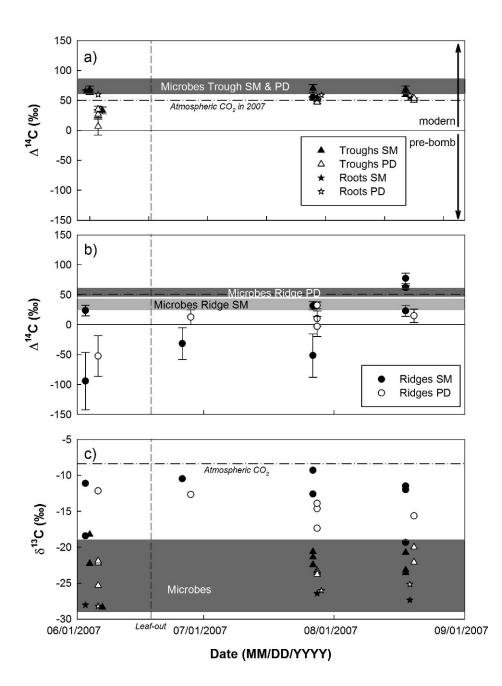


FIGURE 4. Seasonal pattern of the Δ^{14} C signatures of soil, root respiration, and microbial respiration at South Mountain and Polar Desert in (a) troughs and (b) ridges. Soil respiration Δ^{14} C (average \pm SD, n = 3) are corrected for contributions of CO₂ in ambient air based on observed $\delta^{13}C$ ratios (panel c), using either the δ^{13} C of root respiration (-28%) in June, -26% in July and August) or the range of $\delta^{13}C$ of microbial respiration (-19 to -29%) as a proxy for the δ^{13} C of soil respiration. (c) Seasonal pattern of observed $\delta^{13}C$ ratios of soil, root respiration, and microbial respiration at South Mountain and Polar Desert. Symbols represent point measurements of soil or root respiration, horizontal bars indicate observed range of microbial respiration at 0-10 cm soil depth (average \pm SE). Dash-dotted lines indicate $\Delta^{14}C$ or $\delta^{13}C$ ratios of CO_2 in ambient air (50.34 ± 0.65%, -8.37 ± 0.11 ; n = 7), and dashed line day of leaf-out.

microorganisms decomposing relatively young C pools. This pattern has been observed before in northern soils (Wookey et al., 2002.; Hirsch et al., 2003; Czimczik et al., 2006) and demonstrates the difficulty of detecting slow, but potentially sustained decomposition of older soil C pools in surface fluxes.

For spring, when the active layer is shallow, we can estimate the maximum contribution of decomposition of old organic C present at 0–10 cm ($F_{microbes}$) to the observed soil respiration flux (F_{SR}) relative to root respiration (F_{roots}) using an isotope mass balance approach:

$$F_{SR} = F_{roots} + F_{microbes} = 1 \tag{3}$$

For this calculation, we used an average soil respiration flux of 10 or 30 mg C m⁻² hr⁻¹ in ridges or troughs (Figs. 3a, 3b), the ¹⁴C signature of root respiration (Fig. 4a) and the oldest observed ¹⁴C signature of bulk organic matter (Table 2). For soil respiration to reach the observed ¹⁴C signatures between +3 and -106%, the

decomposition of bulk organic matter would have to contribute a maximum of 20–60% of the total flux—depending on the assumed age of the old C pool (Table 3). After leaf-out, the old C source must have contributed less.

Accelerated warming and/or changes in winter and/or summer precipitation—all of which may lead to a deepening of the active layer (Lawrence et al., 2008a; Nowinski et al., 2010)— may result in the release of greater amounts of older soil C into the atmosphere (Schuur et al., 2009). To estimate at what rate older C is being lost, soil C pools need to be better constrained, the actual C sources of the microbial population need to be assessed within the entire active layer, and soils need to be monitored over longer time periods.

Conclusions

Non-sorted, striped Cryosols in Northwest Greenland contain old C pools (Δ^{14} C modern to -470%), located within the top

TABLE 3

Estimated maximum contribution of microbial decomposition of bulk soil organic matter (SOM) to the soil respiration flux at the beginning of the growing season, assuming an active layer depth of <10 cm.

$\frac{\text{Soil respiration}}{\text{Flux (mg C m}^{-2} \text{ hr}^{-1})}$		Roots		SOM decomposition Flux (%)	
		Δ^{14} C (‰)	SOM		
Troughs					
30	3	59	-52	50	
			-82	40	
Ridges South Mountain					
10	-106	59	-332	40	
Ridges Polar Desert					
10	-61	59	-148	60	
			-470	20	

10 cm of mineral soil. Before leaf-out, we estimated that older C pools (fixed by plants prior to 1950) contributed a maximum of 20–60% to the net soil respiration flux. After leaf-out, the contribution of older C pools to net soil respiration declined, but was still apparent in the soil respiration flux from barren ridges as well as in the pore space of mineral soils. Soil respiration in vegetated troughs became dominated by more recently fixed C (between 1950 and 2007). As air temperatures continue to warm in this region of the Arctic, we anticipate a deeper active layer, increased microbial degradation of older soil C pools, and a larger contribution of older C pools to the soil respiration flux.

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